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(21) International Application Number: PCT/US00/00641			(US). HILLMAN, Jennifer, L. [US/US]; 230 Monroe Drive #12, Mountain View, CA 94040 (US). TANG, Y., Tom [US/US]; 4230 Ranwick Court, San Jose, CA 95118 (US). AZIMZAI, Yalda [US/US]; 2045 Rock Springs Drive, Hayward, CA 94545 (US). BAUGHN, Mariah, R. [US/US]; 14244 Santiago Road, San Leandro, CA 94577 (US). LAL, Preeti [US/US]; 2382 Lass Drive, Santa Clara, CA 95054 (US). YUE, Henry [US/US]; 826 Lois Avenue, Sunnyvale, CA 94087 (US). LU, Dyung, Aina, M. [US/US]; 55 Park Belmont Park, San Jose, CA 95136 (US).
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(63) Related by Continuation (CON) or Continuation-in-Part (CIP) to Earlier Applications US 60/172,247 (CIP) Filed on 11 January 1999 (11.01.99) US 60/132,253 (CIP) Filed on 3 May 1999 (03.05.99) US 60/136,653 (CIP) Filed on 27 May 1999 (27.05.99)			(74) Agents: HAMLET-COX, Diana et al.; Incyte Pharmaceuticals, Inc., 3174 Porter Drive, Palo Alto, CA 94304 (US).
(71) Applicant (for all designated States except US): INCYTE PHARMACEUTICALS, INC. [US/US]; 3174 Porter Drive, Palo Alto, CA 94304 (US).			(81) Designated States: AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CU, CZ, DE, DK, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, UA, UG, US, UZ, VN, YU, ZW, ARIPO patent (GH, GM, KE, LS, MW, SD, SL, SZ, TZ, UG, ZW), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG).
(72) Inventors; and (75) Inventors/Applicants (for US only): BANDMAN, Olga [US/US]; 366 Anna Avenue, Mountain View, CA 94043			Published <i>Without international search report and to be republished upon receipt of that report.</i>

(54) Title: HUMAN PEPTIDASES

(57) Abstract

The invention provides human peptidases (HPEP) and polynucleotides which identify and encode HPEP. The invention also provides expression vectors, host cells, antibodies, agonists, and antagonists. The invention also provides methods for diagnosing, treating, or preventing disorders associated with expression of HPEP.

Table 2 (cont.)

SEQ ID NO:	Amino Acid Residues	Potential Phosphorylation Sites	Potential Glycosylation Sites	Signature Sequences, Motifs, and Domains	Homologous Sequences	Analytical Methods
14	703	S20 S68 T120 T135 S331 T383 S562 S606 S607 S631 S674 S698 T31 S95 S115 S173 S355 S490 S562 S650	N318 N434 N445 N670	E1 ubiquitin activating enzyme: K352-H442	E1-like protein ubiquitin activating enzyme) [Pichia pastoris] 94262402	MOTIFS BLAST BLIMPS
15	145	T36 S100 S115 T47	N34	Protease serine hydrolase precursor signal zymogen glycoprotein multigene family: L16-Q64, G87-K140 Trypsin: L25-Q64, S84-N142	Matriptase (serine protease) [Homo sapiens] q5359675, g6002714 Epithrin (membrane bound serine protease) [Mus musculus] g4104970	MOTIFS BLAST BLIMPS
16	518	S74 T252 S151 T169 T245 S312 S361 T419 S462 S502 S16 S70 S98 S133 T301 S331 S428 T516 Y334	N234	Dipeptidyl peptidase IV: H255-L305, E326-Q352, E379-P411	Dipeptidyl peptidase IV [Stenotrophomonas maltophilia] q1753197	MOTIFS BLAST BLIMPS

Pro Thr Ser Leu Gly Leu Val Pro His Gln Ile Arg Gly Phe Leu
 620 625 630
 Ser Arg Phe Asp Asn Val Leu Pro Val Ser Leu Ala Phe Asp Lys
 635 640 645
 Cys Thr Ala Cys Ser Ser Lys Val Leu Asp Gln Tyr Glu Arg Glu
 650 655 660
 Gly Phe Asn Phe Leu Ala Lys Val Phe Asn Ser Ser His Ser Phe
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 680 685 690
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 35 40 45
 Ile Thr Pro Arg Met Met Cys Val Gly Phe Leu Ser Gly Gly Val
 50 55 60
 Asp Ser Cys Gln Val Ala Pro Gly Ala Gly Gly Arg Gln Val Gly
 65 70 75
 Pro Gly Arg Gly Gly Thr Gly Asp Ser Pro Ala Gly Leu Val Ser
 80 85 90
 Ala Gln Gly Asp Ser Gly Gly Pro Leu Ser Ser Val Glu Ala Asp
 95 100 105
 Gly Arg Ile Phe Gln Ala Gly Val Val Ser Trp Gly Asp Gly Cys
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Phe	Gly	Ile	Thr	Ser	Tyr	Asp	Phe	His	Ser	Glu	Ser	Gly	Leu	Phe
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Leu	Phe	Gln	Ala	Ser	Asn	Ser	Leu	Phe	His	Cys	Arg	Asp	Gly	
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Trp	Trp	Cys	Pro	Thr	Ala	Ser	Trp	Glu	Gly	Ser	Glu	Gly	Leu	Lys
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Thr	Leu	Arg	Ile	Leu	Tyr	Glu	Glu	Val	Asp	Glu	Ser	Glu	Val	Glu
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Val	Ile	His	Val	Pro	Ser	Pro	Ala	Leu	Glu	Glu	Arg	Lys	Thr	Asp
				320				325						330
Ser	Tyr	Arg	Tyr	Pro	Arg	Thr	Gly	Ser	Lys	Asn	Pro	Lys	Ile	Ala
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Ser	Thr	Gln	Glu	Lys	Glu	Leu	Val	Gln	Pro	Phe	Ser	Ser	Leu	Phe
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Pro	Lys	Val	Glu	Tyr	Ile	Ala	Arg	Ala	Gly	Trp	Thr	Arg	Asp	Gly
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Lys	Tyr	Ala	Trp	Ala	Met	Phe	Leu	Asp	Arg	Pro	Gln	Gln	Trp	Leu
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 425 430 435
 Val Gln Pro Tyr Val Val Tyr Glu Glu Val Thr Asn Val Trp Ile
 440 445 450
 Asn Val His Asp Ile Phe Tyr Pro Phe Pro Gln Ser Glu Gly Glu
 455 460 465
 Asp Glu Leu Cys Phe Leu Arg Ala Asn Glu Cys Lys Thr Gly Phe
 470 475 480
 Cys His Leu Tyr Lys Val Thr Ala Val Leu Lys Ser Gln Gly Tyr
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 50 55 60
Gly Cys Cys Glu Asp Arg Asp Asp Gly Cys Val Thr Glu Phe Tyr
 65 70 75
Ala Ala Asn Ala Leu Cys Tyr Cys Asp Lys Phe Cys Asp Arg Glu
 80 85 90
Asn Ser Asp Cys Cys Pro Asp Tyr Lys Ser Phe Cys Arg Glu Glu
 95 100 105
Lys Glu Trp Pro Pro His Thr Gln Pro Trp Tyr Pro Glu Gly Cys
 110 115 120
Phe Lys Asp Gly Gln His Tyr Glu Glu Gly Ser Val Ile Lys Glu
 125 130 135
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Gln His Val Cys Leu Val Arg Ser Glu Leu Ile Glu Gln Val Asn
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Lys Gly Asp Tyr Gly Trp Thr Ala Gln Asn Tyr Ser Gln Phe Trp
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Gly Met Thr Leu Glu Asp Gly Phe Lys Phe Arg Leu Gly Thr Leu
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Pro Pro Ser Pro Met Leu Leu Ser Met Asn Glu Met Thr Ala Ser